SEQUENCE LISTING

<110> Joelle THONNARD

<120> Novel Compounds

<130> BM45399

<150> 9918034.1
<151> 1999-07-30

<160> 4

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 336
<212> DNA
<213> Moraxella catarrhalis

<400> 1
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accatcatta agggcaagac caataagcaa gagatttcta gtagatttgg ttctgctgat   180
agcatctctt ttatgatagt ggtaataaaa tttggacata ccgccatact cgctccaaac   240
cgatggcaag aaattttatc ccttataatt tcttttcttt gggtgaaacc atacagacca   300
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<210> 2
<211> 111
<212> PRT
<213> Moraxella catarrhalis

<400> 2
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Leu Leu Val Thr Gly Cys Val Ser Thr Gly Asn Val Ala Met Lys Glu
            20                  25                  30
Gln Asn Gln Gln Thr Ile Glu Gln Thr Ile Ile Lys Gly Lys Thr Asn
        35                  40                  45
Lys Gln Glu Ile Ser Ser Arg Phe Gly Ser Ala Asp Ser Ile Ser Phe
    50                  55                  60
Met Ile Val Val Ile Lys Phe Gly His Thr Ala Ile Leu Ala Pro Asn
65                  70                  75                  80
Arg Trp Gln Glu Ile Leu Ser Leu Ile Ile Ser Phe Leu Trp Val Lys
                85                  90                  95
Pro Tyr Arg Pro Lys Asn Leu Ser Phe Tyr Leu Thr Ala Lys Ala
            100                 105                 110
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<210> 3
<211> 987
<212> DNA

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<213> Moraxella catarrhalis

<400> 3
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cctgtcgttc agcagattga tgacatttgc caagcaccag ccatgcaagc aaatctacaa     180
aatggtatca agcagtccat tcttgatgcc gtgatgagcc aaatgacgga tgcagacccc     240
aaccaacgcc tcgccataca aaataccatt ggtcagcagc tgaatacttt acaaatcacc     300
acccaaaatg ccacgaattt tgccgatagc tgtatggcgg atgtacatat aacggtaaac     360
ccccaagatt tggcgaatgc agaatttgct tttgctcgtt caggggtaac tttattacag     420
cgtgctagcc aagatcaagt tgagttttat aatggcacaa ttgtcgccaa acaaataact     480
tatcaaatgg tcaatggtaa tgtggtgatg tatggcaaca atcataatgc cattcttctg     540
attgccgata ttttggcagc ttctacaagc agtttgccac aagtatccat acaatctgat     600
gttaccgcac gaccacaagc aattgagcga ctgccagagg caccaatagc tatgccatca     660
aatcctcaag aagattctag tgtaacgacg tacattgaac aaaagccagc acctaacgca     720
caggtgtcat ctcgcccacg cagcgagatg tcttcaaata attcagcaca gacacccaca     780
caaaactcgg tagggcaatc ttctgcagct ggtagcaccc ctcgtgtgga tcgtgatagt     840
caagccaaag ccaataccga gcgtaccacc gagcgttctg caaataagac .atctcaagac     900
ttaccacacc cacagccacc aacggcgaat gccagtagtg atggcaaaac aagtatcagt     960
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<210> 4
<211> 328
<212> PRT
<213> Moraxella catarrhalis

<400> 4
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Thr Ala Ala Leu Leu Thr Leu Ala Gly Cys Ser His Asp Asn His Arg
             20                  25                  30
Asp Asp Glu Met Ala Val Ala Glu Pro Val Val Gln Gln Ile Asp Asp
         35                  40                  45
Ile Cys Gln Ala Pro Ala Met Gln Ala Asn Leu Gln Asn Gly Ile Lys
     50                  55                  60
Gln Ser Ile Leu Asp Ala Val Met Ser Gln Met Thr Asp Ala Asp Pro
65                  70                  75                  80
Asn Gln Arg Leu Ala Ile Gln Asn Thr Ile Gly Gln Gln Leu Asn Thr
                 85                  90                  95
Leu Gln Ile Thr Thr Gln Asn Ala Thr Asn Phe Ala Asp Ser Cys Met
             100                 105                 110
Ala Asp Val His Ile Thr Val Asn Pro Gln Asp Leu Ala Asn Ala Glu
         115                 120                 125
Phe Ala Phe Ala Arg Ser Gly Val Thr Leu Leu Gln Arg Ala Ser Gln
     130                 135                 140
Asp Gln Val Glu Phe Tyr Asn Gly Thr Ile Val Ala Lys Gln Ile Thr
145                 150                 155                 160
Tyr Gln Met Val Asn Gly Asn Val Val Met Tyr Gly Asn Asn His Asn
                 165                 170                 175
Ala Ile Leu Leu Ile Ala Asp Ile Leu Ala Ala Ser Thr Ser Ser Leu
             180                 185                 190
Pro Gln Val Ser Ile Gln Ser Asp Val Thr Ala Arg Pro Gln Ala Ile
             195                 200                 205
Glu Arg Leu Pro Glu Ala Pro Ile Ala Met Pro Ser Asn Pro Gln Glu
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            210                215                   220
Asp Ser Ser Val Thr Thr Tyr Ile Glu Gln Lys Pro Ala Pro Asn Ala
225                     230                235                   240
Gln Val Ser Ser Arg Pro Arg Ser Glu Met Ser Ser Asn Asn Ser Ala
                245                     250                255
Gln Thr Pro Thr Gln Asn Ser Val Gly Gln Ser Ser Ala Ala Gly Ser
                260                     265                270
Thr Pro Arg Val Asp Arg Asp Ser Gln Ala Lys Ala Asn Thr Glu Arg
                275                     280                285
Thr Thr Glu Arg Ser Ala Asn Lys Thr Ser Gln Asp Leu Pro His Pro
                290                     295                300
Gln Pro Pro Thr Ala Asn Ala Ser Ser Asp Gly Lys Thr Ser Ile Ser
305                     310                315                   320
Ile Val Glu Ser Asn Glu Thr Tyr
                325
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